Sikandar Amanullah

List of Publications by Year in descending order

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759233 642732 24 570 12 23 h-index g-index citations papers 25 25 25 543 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Volatile terpenoids: multiple functions, biosynthesis, modulation and manipulation by genetic engineering. Planta, 2017, 246, 803-816.	3.2	174
2	Comparative transcriptome analysis of two contrasting watermelon genotypes during fruit development and ripening. BMC Genomics, 2017, 18, 3.	2.8	72
3	Mapping of powdery mildew resistance genes in melon (Cucumis melo L.) by bulked segregant analysis. Scientia Horticulturae, 2017, 220, 160-167.	3.6	59
4	QTL mapping for melon (Cucumis melo L.) fruit traits by assembling and utilization of novel SNPs based CAPS markers. Scientia Horticulturae, 2018, 236, 18-29.	3.6	36
5	Application of nano-titanum dioxide coating on fresh Highbush blueberries shelf life stored under ambient temperature. LWT - Food Science and Technology, 2021, 137, 110422.	5.2	31
6	GsSNAP33, a novel Glycine soja SNAP25-type protein gene: Improvement of plant salt and drought tolerances in transgenic Arabidopsis thaliana. Plant Physiology and Biochemistry, 2017, 119, 9-20.	5.8	20
7	Comparative analysis of single nucleotide polymorphisms in the nuclear, chloroplast, and mitochondrial genomes in identification of phylogenetic association among seven melon (<i>Cucumis melo</i> L.) cultivars. Breeding Science, 2016, 66, 711-719.	1.9	19
8	Genome-Wide Analysis of the Peroxidase Gene Family and Verification of Lignin Synthesis-Related Genes in Watermelon. International Journal of Molecular Sciences, 2022, 23, 642.	4.1	19
9	Genetic linkage mapping and QTLs identification for morphology and fruit quality related traits of melon by SNP based CAPS markers. Scientia Horticulturae, 2021, 278, 109849.	3.6	18
10	Identification of putative genetic regions for watermelon rind hardness and related traits by BSA-seq and QTL mapping. Euphytica, 2021, 217, 1.	1.2	15
11	Detection of putative QTL regions associated with ovary traits in melon using SNP-CAPS markers. Scientia Horticulturae, 2020, 270, 109445.	3.6	14
12	Functional Characterization of Hedychium coronarium J. Koenig MYB132 Confers the Potential Role in Floral Aroma Synthesis. Plants, 2021, 10, 2014.	3.5	14
13	Fine Mapping of Cla015407 Controlling Plant Height in Watermelon. Journal of the American Society for Horticultural Science, 2021, 146, 196-205.	1.0	13
14	Genetic mapping reveals a candidate gene for egusi seed in watermelon. Euphytica, 2019, 215, 1.	1.2	12
15	Mapping of genetic loci controlling fruit linked morphological traits of melon using developed CAPS markers. Molecular Biology Reports, 2022, 49, 5459-5472.	2.3	9
16	Aloe vera Coating Efficiency on Shelf Life of Eggplants at Differential Storage Temperatures. The Journal of Northeast Agricultural University, 2016, 23, 15-25.	0.1	8
17	Identification of QTLs linked with watermelon fruit and seed traits using GBS-based high-resolution genetic mapping. Scientia Horticulturae, 2022, 303, 111237.	3.6	8
18	Exogenous Methyl Jasmonate Application Improved Physio-Biochemical Attributes, Yield, Quality, and Cadmium Tolerance in Fragrant Rice. Frontiers in Plant Science, 2022, 13, 849477.	3.6	7

#	Article	IF	CITATIONS
19	Genetic Mapping and QTL Analysis of Stigma Color in Melon (Cucumis melo L.). Frontiers in Plant Science, 2022, 13, .	3.6	7
20	CAPS marker-base genetic linkage mapping and QTL analysis for watermelon ovary, fruit and seed-related traits. Euphytica, 2022, 218, $1.$	1.2	5
21	Development of Whole Genome SNP-CAPS Markers and Preliminary QTL Mapping of Fruit Pedicel Traits in Watermelon. Frontiers in Plant Science, 2022, 13, .	3.6	5
22	Comparative Transcriptome Analysis Identified Key Pathways and Genes Regulating Differentiated Stigma Color in Melon (Cucumis melo L.). International Journal of Molecular Sciences, 2022, 23, 6721.	4.1	3
23	QTL mapping for significant seed traits of watermelon (Citrullus lanatus Schrad). Pakistan Journal of Botany, 2021, 53, .	0.5	1
24	Molecular cloning, characterization and expression analysis of two 12-oxophytodienoate reductases (NtOPR1 and NtOPR2) from Nicotiana tabacum. Molecular Biology Reports, 2022, 49, 5379-5387.	2.3	1